

010590

#8



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ENTERED

## RAW SEQUENCE LISTING

DATE: 03/19/2002 P.5

PATENT APPLICATION: US/09/883,848A

TIME: 15:13:31

Input Set : A:\CIBT-P01-119 Sequence Listing.txt

Output Set: N:\CRF3\03192002\I883848A.raw

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3 <110> APPLICANT: Ling, L.
4   Sanicola-Nadel, M.
6 <120> TITLE OF INVENTION: ANGIOGENESIS-MODULATING COMPOSITIONS AND USES
8 <130> FILE REFERENCE: CIBT-P01-119
10 <140> CURRENT APPLICATION NUMBER: 09/883,848A
11 <141> CURRENT FILING DATE: 2001-06-18
13 <150> PRIOR APPLICATION NUMBER: 60/211,919
14 <151> PRIOR FILING DATE: 2000-06-16
16 <160> NUMBER OF SEQ ID NOS: 48
18 <170> SOFTWARE: PatentIn Ver. 2.1
20 <210> SEQ ID NO: 1
21 <211> LENGTH: 1277
22 <212> TYPE: DNA
23 <213> ORGANISM: Gallus gallus
25 <400> SEQUENCE: 1
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28 aagctgaccc cgtagccta taagcagttt attcccaatg tggcagagaa gaccctaggg 180
29 gccagtggaa gatatgaagg gaagatcaca agaaactccg agagatttaa agaactaacc 240
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31 atgactcagc gctgcaagga caagctgaat gccctggcga tctcggtgat gaaccagtgg 360
32 cccgggggtga agctgagggt gaccgagggc tgggacgagg atggccatca ctccgaggaa 420
33 tcgctgcact acgagggtcg cgccgtggac atcaccacgt cggatcggga ccgcagcaag 480
34 tacggaatgc tggcccgccct cgccgtcgag gccggcttcg actgggtcta ctacgagtcc 540
35 aaggcgcaca tccactgctc cgtaaaagca gaaaactcag tggcagcgaa atcaggaggc 600
36 tgcttccctg gctcagccac agtgacactg gagcatggag gcaccaagct ggtgaaggac 660
37 ctgagccctg gggaccgcgt gctggctgct gacgcggacg gccggtgct ctacagtgc 720
38 ttctcacct tcctcgaccg gatggacagc tcccgaagc tcttctacgt catcgagacg 780
39 cggcagcccc gggcccggt gctactgacg gcggcccacc tgctctttgt ggccccccag 840
40 cacaaccagt cggaggccac aggtgccacc agtggccagg cgctcttcgc cagcaacgtg 900
41 aagcctggcc aacgtgtcta tgtgtgggc gagggcgggc agcagctgct gccggcgtct 960
42 gtccacagcg tctcattgcg ggaggaggcg tccggagcct acgccccact caccgcccag 1020
43 ggcaccatcc tcatcaaccg ggtgttggcc tcctgctacg ccgtcatcga ggagcacagt 1080
44 tgggcccatt gggccttcgc accattccgc ttggctcagg ggctgctggc cgccctctgc 1140
45 ccagatgggg ccatccctac tgccgccacc accaccactg gcatccattg gtactcacgg 1200
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47 gtggcaccgg ccagctg                                     1277
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55 <400> SEQUENCE: 2
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57 cagagctgcg ggccggggcg aggaccggtt ggccggcgcg gttatgtgcg caagcaactt 120
58 gtgcctctgc tatacaagca gtttgtgccc agtatgcccg agcggaccct gggcgcgagt 180
59 gggccagcgg aggggagggg aacaaggggg tcggagcgct tccgggacct cgtacccaac 240
60 tacaaccccc acataatctt caaggatgag gagaacagcg gcgcagaccg cctgatgaca 300
61 gagcgttgca aagagcgggg gaacgctcta gccatcgcgg tgatgaacat gtggcccggg 360
62 gtacgcctac gtgtgactga aggctgggac gaggacggcc accacgcaca ggattcactc 420
63 cactacgaag gccgtgcctt ggacatcacc acgtctgacc gtgaccgtaa taagtatggg 480
64 ttgttgggcg gcctagctgt ggaagccgga ttcgactggg tctactacga gtcccgaac 540
65 cacatccacg tatcgtgcaa agctgataac tctactggcg tccgagccgg aggtgcttt 600
66 ccgggaaatg ccacggtgcg cttgcgagc ggcgaaacgga aggggctgag ggaactacat 660
67 cgtggtgact gggtaactgg cgctgatgca gcgggcccag tggtaaccac gccagtgtg 720
68 ctcttcctgg accgggatct gcacgcccgc cctcgttcg tggctgtgga gaccgagcg 780
69 cctccgcgca aactgttgct cacaccctgg catctggtgt tcgctgctcg cgggccagcg 840
70 cctgctccag gtgactttgc accggtgttc gcgcgcgct tacgtgctgg cgactcgtgtg 900
71 ctggctcccg gcggggacgc gctccagccg gcgcgcgtag cccgcgtggc gcgcgaggaa 960
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73 gcctcctgct acgcggttct agagagtcac cagtggggcc accgcgcctt cggccctttg 1080
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86 cctcgcaagc tcgtgcctct tgctacaag cagttcagcc ccaacgtgcc ggagaagacc 180
87 ctgggcgcca gcgggcgcta cgaaggcaag atcgcgcgca gctctgagcg cttcaaagag 240
88 ctcaccccca actacaatcc cgacatcatc ttcaaggacg aggagaacac gggtgccgac 300
89 cgcctcatga cccagcgctg caaggaccgt ctgaactcac tggccatctc tgtcatgaac 360
90 cagtggcctg gtgtgaaaact gcgggtgacc gaaggccggg atgaagatgg ccatcactca 420
91 gaggagtctt tacactatga gggccgcgcg gtggatatca ccacctcaga ccgtgaccga 480
92 aataagtatg gactgctggc gcgcttagca gtggaggccg gcttcgactg ggtgtattac 540
93 gagtccaagg cccacgtgca ttgctctgtc aagtctgagc attcggccgc tgccaagaca 600
94 ggtggctgct ttcttgccgg agcccagggt cgcttagaga acggggagcg tgtggccctg 660
95 tcagctgtaa agccaggaga ccgggtgctg gccatggggg aggatgggac cccaccttc 720
96 agtgatgtgc ttattttcct ggaccgcgag ccaaaccggc tgagagcttt ccaggtcatc 780
97 gagactcagg atcctccgcg tcggctggcg ctacgcctg cccacctgct cttcattgcg 840
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99 ggccaatatg tgctggtatc aggggtacca ggccctcagc ctgctcgggt ggcagctgtc 960
100 tccacccacg tggcccttgg gtcctatgct cctctcaca ggcatgggac acttgtgggtg 1020
101 gaggatgttg tggcctcctg ctttgcagct gtggtgacc accatctggc tcagttggcc 1080
102 ttctggcccc tgcgactgtt tcccagtttg gcatggggca gctggacccc aagtgagggt 1140
103 gttcactcct accctcagat gctctaccgc ctggggcgct tcttgctaga agagagcacc 1200
104 ttccatccac tgggcatgtc tggggcagga agctgaaggg actctaacca ctgccctcct 1260
105 ggaactgctg tgcggtgata c
108 <210> SEQ ID NO: 4
109 <211> LENGTH: 1313
110 <212> TYPE: DNA

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Output Set: N:\CRF3\03192002\I883848A.raw

111 &lt;213&gt; ORGANISM: Mus musculus

113 &lt;400&gt; SEQUENCE: 4

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116 acccttttag cctacaagca gtttattccc aacgtagccg agaagaccct agggggccagc 180
117 ggcagatatg aagggaagat cacaagaaac tccgaacgat ttaaggaact caccctccaat 240
118 tacaaccccg acatcatatt taaggatgag gaaaacacgg gagcagaccg gctgatgact 300
119 cagaggtgca aagacaagtt aaatgccttg gccatctctg tgatgaacca gtggcctgga 360
120 gtgaggctgc gactgaccga gggctgggat gaggacggcc atcattcaga ggagtctcta 420
121 cactatgagg gtcgagcagt ggacatcacc acgtccgacc gggaccgcag caagtacggc 480
122 atgctggctc gcctggctgt ggaagcaggt ttcgactggg tctactatga atccaaagct 540
123 cacatccact gttctgtgaa agcagagaaac tccgtggcgg ccaaattccg cggtgttttc 600
124 ccgggatccg ccaccgtgca cctggagcag ggcggcacca agctggtgaa ggacttacgt 660
125 cccggagacc gcgtgctggc ggctgacgac cagggccggc tgctgtacag cgacttcctc 720
126 accttcctgg accgcgacga aggcgccaag aaggtcttct acgtgatcga gacgtggag 780
127 ccgcgcgagc gcctgctgct caccgcgcg cactgtctct tcgtggcgcc gcacaacgac 840
128 tcggggccca cggccgggccc aagcgcgctc tttgccagcc gcgtgcgccc cgggcagcgc 900
129 gtgtacgtgg tggctgaacg cggcggggac cgccggctgc tgcccgccg ggtgcacagc 960
130 gtgacgctgc gagaggagga ggcggggcgc tacgcgccgc tcacggcgca cggcaccatt 1020
131 ctcatcaacc ggggtgctcg ctcgtgctac gctgtcatcg aggagcacag ctgggcacac 1080
132 cgggccttcg cgcctttccg cctggcgcac gcgctgctgg ccgcgctggc acccgcccgc 1140
133 acggacggcg ggggcggggg cagcatccct gcagcgcaat ctgcaacgga agcgaggggc 1200
134 gcggagccga ctgcgggcat ccactggtac tcgcagctgc tctaccacat tggcacctgg 1260
135 ctgttggaac gcgagaccat gcatcccttg ggaatggcgg tcaagtccag ctg 1313

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138 &lt;210&gt; SEQ ID NO: 5

139 &lt;211&gt; LENGTH: 1256

140 &lt;212&gt; TYPE: DNA

141 &lt;213&gt; ORGANISM: Brachydanio rerio

143 &lt;400&gt; SEQUENCE: 5

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144 atgcggcttt tgacgagagt gctgctggtg tctcttctca ctctgtcctt ggtggtgtcc 60
145 ggactggcct gcggtcctgg cagaggctac ggcagaagaa gacatccgaa gaagctgaca 120
146 cctctcgctt acaagcagtt cataccta atgtcgcgga agaccttagg ggccagcggc 180
147 agatacgagg gcaagataac gcgcaattcg gagagattta aagaacttac tccaaattac 240
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152 ctgtctcgcc tagctgttga ggctggattt gactgggtct attacgagtc caaagcccac 540
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154 ggttcggctc tggctcgcct ccaggacgga ggacagaagg ccgtgaagga cctgaacccc 660
155 ggagacaagg tgctggcggc agacagcgcg ggaaacctgg tggtcagcga cttcatcatg 720
156 ttacagacc gagactccac gacgcgacgt gtgttttacg tcatagaaac gcaagaaccc 780
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159 gttgttgatg atagcggta cgttaaatct gtcacgtgac agcggatata cagggaggag 960
160 cagcggggct cgttcgcacc agtgactgca catgggacca ttgtggtcga cagaatactg 1020
161 gcgtcctgtt acgccgtaat agaggaccag ggccttgccg atttggcctt cgcgcccgc 1080
162 aggtcttatt attacgtgtc atcattcctg tccccaaaa ctccagcagt cgggtccaatg 1140
163 cgactttaca acaggagggg gtccactggt actccaggct cctgtcatca aatgggaacg 1200

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164 tggcttttgg acagcaacat gcttcacatcct ttgggggatgt cagtaaactc aagctg      1256
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167 <211> LENGTH: 1425
168 <212> TYPE: DNA
169 <213> ORGANISM: Homo sapiens
171 <220> FEATURE:
172 <221> NAME/KEY: modified_base
173 <222> LOCATION: (1387...1389)
174 <223> OTHER INFORMATION: n=a, c, g, or t
176 <400> SEQUENCE: 6
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178 ggactggcgt gcggaccggg caggggggttc gggaagagga ggcaccccaa aaagctgacc 120
179 ccttttagcct acaagcagtt tatccccaat gtggccgaga agaccctagg cgccagcgga 180
180 aggtatgaag ggaagatctc cagaaaactcc gagcgattta aggaactcac cccaattac 240
181 aaccccgaca tcatatttaa ggatgaagaa aacaccggag cggacaggct gatgactcag 300
182 aggtgtaagg acaagttgaa cgctttggcc atctcgtgta tgaaccagtg gccaggagtg 360
183 aaactgcggg tgaccgaggg ctgggacgaa gatggccacc actcagagga gtctctgcac 420
184 tacgagggcc gcgcagtgga catcaccacg tctgaccgcg accgcagcaa gtacggcatg 480
185 ctggcccgcc tggcgggtgga ggccggcttc gactgggtgt actacgagtc caaggcacat 540
186 atccactgct cggtgaaagc agagaactcg gtggcggcca aatcgggagg ctgcttcccg 600
187 ggctcggcca cggtgacact ggagcagggc ggacccaagc tggtgaaagga cctgagcccc 660
188 ggggaccgcg tgctggcggc ggacgaccag ggccggctgc tctacagcga ctctctcact 720
189 ttcttgacc gcgacgacgg cgccaagaag gtctttctac tgatcgagac gcgggagccg 780
190 cgcgagcgcc tgctgctcac cgccgcgcac ctgctctttg tggcgccgca caacgactcg 840
191 gccaccgggg agcccagggc gtccctcgggc tcggggccgc cttccggggg cgcactgggg 900
192 cctcggggcg tgttcgccag ccgcgtgcgc ccggggccagc gcgtgtacgt ggtggccgag 960
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194 gccgcgggcg cctacgcgcc gctcacggcc cagggcacca ttctcatcaa ccgggtgctg 1080
195 gcctcgtgct acgcggtcat cgaggagcac agctgggcgc accgggcctt cgcgcccttc 1140
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197 agcggcgggc gggaccgcgg gggcgggcggc ggcagagtag ccctaaccgc tccagggtgt 1260
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199 caaataggca cctggctcct ggacagcgag gccctgcacc cgctgggcat ggcggtcaag 1380
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204 <211> LENGTH: 1622
205 <212> TYPE: DNA
206 <213> ORGANISM: Homo sapiens
208 <400> SEQUENCE: 7
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210 cccggctccg gccccgactg caattctgcc tggctcgtgt gctgctgctg gtggtgcccg 120
211 cggcatgggg ctgcgggccc ggtcgggttg tgggcagccg ccggcgaccg ccacgcaaac 180
212 tcgtgccgct cgctacaag cagttcagcc ccaatgtgcc cgagaagacc ctgggcgcca 240
213 gcggagccta tgaaggcaag atcgctcgca gctccgagcg cttcaaggag ctacccccca 300
214 attacaatcc agacatcac ttcaaggacg aggagaacac aggcgcccag cgccctcatg 360
215 cccagcgctg caaggaccgc ctgaactcgc tggctatctc ggtgatgaac cagtggcccc 420
216 gtgtgaagct gcgggtgacc gagggctggg acgaggacgg ccaccactca gaggagtccc 480
217 tgcatatga gggccgcgcg gtggacatca ccacatcaga ccgcgaccgc aataagtatg 540
218 gactgctggc gcgcttgga gtggaggccg gctttgactg ggtgtattac gagtcaaaag 600

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219 cccacgtgca ttgctccgtc aagtccgagc actcggccgc agccaagacg ggcggctgct 660
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221 ggccggggaga ccgtgtgctg gccatggggg aggatgggag cccacacctc agcgtatgct 780
222 tcattttcct ggaccgcgag cccacaggc tgagagcctt ccaggtcacg gagactcagg 840
223 accccccaag ccgcctggca ctcacaccg ctacactgct ctttacggct gacaatcaca 900
224 cggagccggc agcccgttc cgggccacat ttgccagcca cgtgcagcct ggccagtacg 960
225 tgcgtgtggc tgggttgcca ggctgcagc ctgcccgcgt ggcatgtgc tctacacacg 1020
226 tggccctcgg ggctacgcc ccgtcacaa agcatgggac actggtgtg gaggatgtgg 1080
227 tggcatcctg ctgcgcggc gtggtgacc accactggc tcagttggc ttctggcccc 1140
228 tgagactcct tcacagcttg gcatggggc gctggacccc gggggagggt gtgcatttgt 1200
229 acccccagct gctctaccg ctggggcgct tctgtctaga agagggcagc ttccaccac 1260
230 tgggcatgtc cggggcaggg agctgaaagg actccaccg tgcctcctg gaactgctgt 1320
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232 ggacactggc tctgccatc tctctgcc tgaagataca ccattgagac ttgactgggc 1440
233 aacaccagcg tccccaccc gcgtcgtgt gtagtcatag agctgcaagc tgagctggcg 1500
234 aggggatgtg tgttgacccc tctctcctag agaccttgag gctggcacgg cgactcccaa 1560
235 ctacgcctgc tctactacg agttttcata ctctgcctcc cccattggga gggccattc 1620
236 cc 1622

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239 &lt;210&gt; SEQ ID NO: 8

240 &lt;211&gt; LENGTH: 1191

241 &lt;212&gt; TYPE: DNA

242 &lt;213&gt; ORGANISM: Homo sapiens

244 &lt;400&gt; SEQUENCE: 8

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246 cagagctgcg ggccggggcg ggggcccgtt ggccggcgcc gctatgcgcg caagcagctc 120
247 gtgccgctac tctacaagca atttgtgccc ggctgtccag agcggaccct gggcgccagt 180
248 gggccagcgg aggggagggt ggcaaggggc tccgagcgt tccgggacct cgtgcccaac 240
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250 gagcgttgca aggagagggt gaacgctttg gccattgccc tgatgaacat gtggcccggg 360
251 gtgcgcctac gactgactga gggctggggc gaggacggcc accacgctca ggattcactc 420
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254 cagctccacg tgcgtgcaa agctgataac tcaactggcg tccgggcccg cggtgcttt 600
255 ccgggaaatg caactgtgcg cctgtggagc ggcgagcgga aagggtgcg ggaactgcac 660
256 cgcggagact gggttttggc ggccgatgcg tcaggccggg tgggtgccac gccgtgctg 720
257 ctcttcctgg accgggactt gcagcgccgg gcttcatttg tggctgtgga gaccgagtgg 780
258 cctccacgca aactgttgc cagccctgg cactggtgt ttgccgctcg agggccggcg 840
259 ccgcgcgccg gcgactttgc accggtgttc gcgcgcggc tacgcgctgg ggactcgggtg 900
260 ctggcgcccg gcggggatgc gcttcggcca gcgcgcgtgg cccgtgtggc gcgggaggaa 960
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263 agactgctgc acgcgctagg ggcgctgct cccggcgggg ccgtccagcc gactggcatg 1140
264 cattggtact ctggctcct ctaccgctta gcggaggagc tactgggctg a 1191

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267 &lt;210&gt; SEQ ID NO: 9

268 &lt;211&gt; LENGTH: 1251

269 &lt;212&gt; TYPE: DNA

270 &lt;213&gt; ORGANISM: Brachydanio rerio

272 &lt;400&gt; SEQUENCE: 9

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273 atggacgtaa ggctgcact gaagcaattt gctttactgt gttttatcag cttgcttctg 60

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Use of n and / or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to ensure a corresponding explanation is present in the <220> to <223> fields of each sequence using n or Xaa.

## VERIFICATION SUMMARY

DATE: 03/19/2002

PATENT APPLICATION: US/09/883,848A

TIME: 15:13:32

Input Set : A:\CIBT-P01-119 Sequence Listing.txt

Output Set: N:\CRF3\03192002\I883848A.raw

L:200 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6  
L:826 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15  
L:1706 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26  
L:1709 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26  
L:1712 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26  
L:1715 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26  
L:1719 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26  
L:1722 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26  
L:1728 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26  
L:1731 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26  
L:1734 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26  
L:1737 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26